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<110> Ono, Yuichi
        Nakagawa, Yasuko
        Sakamoto, Yoshimasa

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Gly Phe Leu Ser Gly Ser Lys Phe Gln Ala Pro Gly Ser Trp Lys Asp
65   70   75   80

Cys Phe Gly Ala Pro Pro Ala Pro Asp Val Leu Arg Ala Asp Arg Ser
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Val Gly Glu Gly Cys Pro Gln Lys Leu Val Thr Ala Asn Leu Leu Arg
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8

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9

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Cys Tyr Ile Thr Gly Trp Gly His Met Gly Asn Lys Met Pro Phe Lys
            995   1000   1005

Leu Gln Glu Gly Glu Val Arg Ile Ile Pro Leu Glu Gln Cys Gln
    1010   1015   1020

Ser Tyr Phe Asp Met Lys Thr Ile Thr Asn Arg Met Ile Cys Ala
    1025   1030   1035

Gly Tyr Glu Ser Gly Thr Val Asp Ser Cys Met Gly Asp Ser Gly
    1040   1045   1050

Gly Pro Leu Val Cys Glu Arg Pro Gly Gly Gln Trp Thr Leu Phe
    1055   1060   1065

Gly Leu Thr Ser Trp Gly Ser Val Cys Phe Ser Lys Val Leu Gly
    1070   1075   1080

Pro Gly Val Tyr Ser Asn Val Ser Tyr Phe Val Gly Trp Ile Glu
    1085   1090   1095

Arg Gln Ile Tyr Ile Gln Thr Phe Leu Gln Lys Lys Ser Gln Gly
    1100   1105   1110

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<211>   1042
<212>   PRT
<213>   Homo sapiens

<400>   4
Met Lys Gln Ser Pro Ala Leu Ala Pro Glu Glu Arg Tyr Arg Arg Ala
1   5   10   15

Gly Ser Pro Lys Pro Val Leu Arg Ala Asp Asp Asn Asn Met Gly Asn
            20   25   30

Gly Cys Ser Gln Lys Leu Ala Thr Ala Asn Leu Leu Arg Phe Leu Leu
        35   40   45

Leu Val Leu Ile Pro Cys Ile Cys Ala Leu Val Leu Leu Leu Val Ile
    50   55   60

Leu Leu Ser Tyr Val Gly Thr Leu Gln Lys Val Tyr Phe Lys Ser Asn
65   70   75   80

Gly Ser Glu Pro Leu Val Thr Asp Gly Glu Ile Gln Gly Ser Asp Val

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                    85                      90                      95
Ile Leu Thr Asn Thr Ile Tyr Asn Gln Ser Thr Val Val Ser Thr Ala

                    100                     105                     110
His Pro Asp Gln His Val Pro Ala Trp Thr Thr Asp Ala Ser Leu Pro
        115                     120                     125

Gly Asp Gln Ser His Arg Asn Thr Ser Ala Cys Met Asn Ile Thr His
        130                     135                     140

Ser Gln Cys Gln Met Leu Pro Tyr His Ala Thr Leu Thr Pro Leu Leu
145                     150                     155                     160

Ser Val Val Arg Asn Met Glu Met Glu Lys Phe Leu Lys Phe Phe Thr
                165                     170                     175

Tyr Leu His Arg Leu Ser Cys Tyr Gln His Ile Met Leu Phe Gly Cys
            180                     185                     190

Thr Leu Ala Phe Pro Glu Cys Ile Ile Asp Gly Asp Asp Ser His Gly
        195                     200                     205

Leu Leu Pro Cys Arg Ser Phe Cys Glu Ala Ala Lys Glu Gly Cys Glu
        210                     215                     220

Ser Val Leu Gly Met Val Asn Tyr Ser Trp Pro Asp Phe Leu Arg Cys
225                     230                     235                     240

Ser Gln Phe Arg Asn Gln Thr Glu Ser Ser Asn Val Ser Arg Ile Cys
                245                     250                     255

Phe Ser Pro Gln Gln Glu Asn Gly Lys Gln Leu Leu Cys Gly Arg Gly
            260                     265                     270

Glu Asn Phe Leu Cys Ala Ser Gly Ile Cys Ile Pro Gly Lys Leu Gln
            275                     280                     285

Cys Asn Gly Tyr Asn Asp Cys Asp Asp Trp Ser Asp Glu Ala His Cys
        290                     295                     300

Asn Cys Ser Glu Asn Leu Phe His Cys His Thr Gly Lys Cys Leu Asn
305                     310                     315                     320

Tyr Ser Leu Val Cys Asp Gly Tyr Asp Asp Cys Gly Asp Leu Ser Asp
                325                     330                     335

Glu Gln Asn Cys Asp Cys Asn Pro Thr Thr Glu His Arg Cys Gly Asp
            340                     345                     350

Gly Arg Cys Ile Ala Met Glu Trp Val Cys Asp Gly Asp His Asp Cys
            355                     360                     365

Val Asp Lys Ser Asp Glu Val Asn Cys Ser Cys His Ser Gln Gly Leu
        370                     375                     380

Val Glu Cys Arg Asn Gly Gln Cys Ile Pro Ser Thr Phe Gln Cys Asp
385                     390                     395                     400

Gly Asp Glu Asp Cys Lys Asp Gly Ser Asp Glu Glu Asn Cys Ser Val
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Ile Gln Thr Ser Cys Gln Glu Gly Asp Gln Arg Cys Leu Tyr Asn Pro
            420   425   430

Cys Leu Asp Ser Cys Gly Gly Ser Ser Leu Cys Asp Pro Asn Asn Ser
            435   440   445

Leu Asn Asn Cys Ser Gln Cys Glu Pro Ile Thr Leu Glu Leu Cys Met
465   455   460

Asn Leu Pro Tyr Asn Ser Thr Ser Tyr Pro Asn Tyr Phe Gly His Arg
465   470   475   480

Thr Gln Lys Glu Ala Ser Ile Ser Trp Glu Ser Ser Leu Phe Pro Ala
            485   490   495

Leu Val Gln Thr Asn Cys Tyr Lys Tyr Leu Met Phe Phe Ser Cys Thr
            500   505   510

Ile Leu Val Pro Lys Cys Asp Val Asn Thr Gly Glu Arg Ile Pro Pro
            515   520   525

Cys Arg Ala Leu Cys Glu His Ser Lys Glu Arg Cys Glu Ser Val Leu
            530   535   540

Gly Ile Val Gly Leu Gln Trp Pro Glu Asp Thr Asp Cys Ser Gln Phe
545   550   555   560

Pro Glu Glu Asn Ser Asp Asn Gln Thr Cys Leu Met Pro Asp Glu Tyr
                565   570   575

Val Glu Glu Cys Ser Pro Ser His Phe Lys Cys Arg Ser Gly Gln Cys
            580   585   590

Val Leu Ala Ser Arg Arg Cys Asp Gly Gln Ala Asp Cys Asp Asp Asp
            595   600   605

Ser Asp Glu Glu Asn Cys Gly Cys Lys Glu Arg Asp Leu Trp Glu Cys
    610   615   620

Pro Ser Asn Lys Gln Cys Leu Lys His Thr Val Ile Cys Asp Gly Phe
625   630   635   640

Pro Asp Cys Pro Asp Tyr Met Asp Glu Lys Asn Cys Ser Phe Cys Gln
            645   650   655

Asp Asp Glu Leu Glu Cys Ala Asn His Ala Cys Val Ser Arg Asp Leu
            660   665   670

Trp Cys Asp Gly Glu Ala Asp Cys Ser Asp Ser Ser Asp Glu Trp Asp
            675   680   685

Cys Val Thr Leu Ser Ile Asn Val Asn Ser Ser Ser Phe Leu Met Val
            690   695   700

His Arg Ala Ala Thr Glu His His Val Cys Ala Asp Gly Trp Gln Glu
705   710   715   720

Ile Leu Ser Gln Leu Ala Cys Lys Gln Met Gly Leu Gly Glu Pro Ser
            725   730   735

```
Val Thr Lys Leu Ile Gln Glu Gln Glu Lys Glu Pro Arg Trp Leu Thr
        740                 745             750

Leu His Ser Asn Trp Glu Ser Leu Asn Gly Thr Thr Leu His Glu Leu
        755             760             765

Leu Val Asn Gly Gln Ser Cys Glu Ser Arg Ser Lys Ile Ser Leu Leu
    770             775             780

Cys Thr Lys Gln Asp Cys Gly Arg Arg Pro Ala Ala Arg Met Asn Lys
785             790             795                 800

Arg Ile Leu Gly Gly Arg Thr Ser Arg Pro Gly Arg Trp Pro Trp Gln
                805             810             815

Cys Ser Leu Gln Ser Glu Pro Ser Gly His Ile Cys Gly Cys Val Leu
            820             825             830

Ile Ala Lys Lys Trp Val Leu Thr Val Ala His Cys Phe Glu Gly Arg
        835             840             845

Glu Asn Ala Ala Val Trp Lys Val Val Leu Gly Ile Asn Asn Leu Asp
    850             855             860

His Pro Ser Val Phe Met Gln Thr Arg Phe Val Lys Thr Ile Ile Leu
865             870             875                 880

His Pro Arg Tyr Ser Arg Ala Val Val Asp Tyr Asp Ile Ser Ile Val
            885             890             895

Glu Leu Ser Glu Asp Ile Ser Glu Thr Gly Tyr Val Arg Pro Val Cys
            900             905             910

Leu Pro Asn Pro Glu Gln Trp Leu Glu Pro Asp Thr Tyr Cys Tyr Ile
        915             920             925

Thr Gly Trp Gly His Met Gly Asn Lys Met Pro Phe Lys Leu Gln Glu
    930             935             940

Gly Glu Val Arg Ile Ile Ser Leu Glu His Cys Gln Ser Tyr Phe Asp
945             950             955                 960

Met Lys Thr Ile Thr Thr Arg Met Ile Cys Ala Gly Tyr Glu Ser Gly
            965             970             975

Thr Val Asp Ser Cys Met Gly Asp Ser Gly Gly Pro Leu Val Cys Glu
            980             985             990

Lys Pro Gly Gly Arg Trp Thr Leu Phe Gly Leu Thr Ser Trp Gly Ser
        995             1000            1005

Val Cys Phe Ser Lys Val Leu Gly Pro Gly Val Tyr Ser Asn Val Ser
    1010            1015            1020

Tyr Phe Val Glu Trp Ile Lys Arg Gln Ile Tyr Ile Gln Thr Phe Leu
1025            1030            1035            1040

Leu Asn
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13

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<210> 5
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:adapter for
      cDNA amplification

<400> 5
cagctccaca acctacatca ttccgt                                    26


<210> 6
<211> 12
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:adapter for
      cDNA amplification

<400> 6
acggaatgat gt                                                   12


<210> 7
<211> 26
<212> DNA
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<220>
<223> Description of Artificial Sequence:adapter for
      cDNA amplification

<400> 7
gtccatcttc tctctgagac tctggt                                    26


<210> 8
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:adapter for
      cDNA amplification

<400> 8
accagagtct ca                                                   12


<210> 9
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:adapter for
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            cDNA amplification

<400> 9
ctgatgggtg tcttctgtga gtgtgt                                    26



<210> 10
<211> 12
<212> DNA
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<220>
<223> Description of Artificial Sequence:adapter for
      cDNA amplification

<400> 10
acacactcac ag                                                   12



<210> 11
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:adapter for
      cDNA amplification

<400> 11
ccagcatcga gaatcagtgt gacagt                                    26



<210> 12
<211> 12
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:adapter for
      cDNA amplification

<400> 12
actgtcacac tg                                                   12



<210> 13
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:adapter for
      cDNA amplification

<400> 13
gtcgatgaac ttcgactgtc gatcgt                                    26



<210> 14
<211> 12
<212> DNA
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:adapter for
     cDNA amplification

<400> 14
acgatcgaca gt   12